

## **STIC Biotechnology Systems Branch**

### **RAW SEQUENCE LISTING** **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/537,847  
Source: Pu/10  
Date Processed by STIC: 2/2/06

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

**FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:**

**<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>**

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)**
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450**
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):**  
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05

## Raw Sequence Listing Error Summary

### ERROR DETECTED

### SUGGESTED CORRECTION

SERIAL NUMBER:

10/537,847

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1. Wrapped Nucleics  
Wrapped Aminos      The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."
  
2. Invalid Line Length      The rules require that a line **not exceed** 72 characters in length. This includes white spaces.
  
3. Misaligned Amino  
Numbering      The numbering under each 5<sup>th</sup> amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.
  
4. Non-ASCII      The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. Please **ensure your subsequent submission is saved in ASCII text**.
  
5. Variable Length      Sequence(s)        contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
  
6. PatentIn 2.0  
"bug"      A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s)       . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**
  
7. Skipped Sequences  
(OLD RULES)      Sequence(s)        missing. If intentional, please insert the following lines for **each** skipped sequence:  
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
 This sequence is intentionally skipped  
  
 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.
  
8. Skipped Sequences  
(NEW RULES)      Sequence(s)        missing. If **intentional**, please insert the following lines for **each** skipped sequence.  
 <210> sequence id number  
 <400> sequence id number  
 000
  
9. Use of n's or Xaa's  
(NEW RULES)      Use of n's and/or Xaa's have been detected in the Sequence Listing.  
 Per 1.823 of Sequence Rules, use of <220>-<223> is **MANDATORY** if n's or Xaa's are present.  
 In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.
  
10. Invalid <213>  
Response      Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence
  
11. Use of <220>      Sequence(s)        missing the <220> "Feature" and associated numeric identifiers and responses.  
 Use of <220> to <223> is **MANDATORY** if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
  
12. PatentIn 2.0  
"bug"      Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
  
13. Misuse of n/Xaa      "n" can **only** represent a single nucleotide; "Xaa" can **only** represent a single amino acid



PCT

## RAW SEQUENCE LISTING

DATE: 02/02/2006

PATENT APPLICATION: US/10/537,847

TIME: 11:46:47

Input Set : A:\SEQLIST.TXT

Output Set: N:\CRF4\02022006\J537847.raw

4 <110> APPLICANT: LIAU, Gene  
 5 STEFANSSON, Steingrimur  
 6 SU, Joseph  
 8 <120> TITLE OF INVENTION: ENDOTHELIAL CELL SPECIFICALLY BINDING  
 9 PEPTIDES  
 11 <130> FILE REFERENCE: 4-32422A/GTI/USN  
 13 <140> CURRENT APPLICATION NUMBER: 10/537,847  
 C--> 14 <141> CURRENT FILING DATE: 2005-06-08  
 16 <150> PRIOR APPLICATION NUMBER: PCT/EP03/014407  
 17 <151> PRIOR FILING DATE: 2003-12-17  
 19 <150> PRIOR APPLICATION NUMBER: 60/434,258  
 20 <151> PRIOR FILING DATE: 2002-12-18  
 22 <160> NUMBER OF SEQ ID NOS: 43  
 24 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
 26 <210> SEQ ID NO: 1  
 27 <211> LENGTH: 9  
 28 <212> TYPE: PRT  
 29 <213> ORGANISM: Artificial Sequence  
 31 <220> FEATURE:  
 32 <223> OTHER INFORMATION: (Artificial peptide sequence)  
 W--> 34 <221> NAME/KEY: PEPTIDE  
 35 <222> LOCATION: (1)...(9)  
 36 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding  
 W--> 38 <400> 1  
 39 Cys Pro Asp Leu His His Met Cys  
 40 1 5  
 43 <210> SEQ ID NO: 2  
 44 <211> LENGTH: 9  
 45 <212> TYPE: PRT  
 46 <213> ORGANISM: Artificial Sequence  
 48 <220> FEATURE:  
 49 <223> OTHER INFORMATION: Artificial peptide sequence  
 W--> 51 <221> NAME/KEY: PEPTIDE  
 52 <222> LOCATION: (1)...(9)  
 53 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding  
 W--> 55 <400> 2  
 56 Cys Leu Gly Gln His Ala Phe Thr Cys  
 57 1 5  
 60 <210> SEQ ID NO: 3  
 61 <211> LENGTH: 9  
 62 <212> TYPE: PRT  
 63 <213> ORGANISM: Artificial Sequence  
 65 <220> FEATURE:

ppt-6  
 Does Not Comply  
 Corrected Diskette Needed

insufficient explanation  
 (give source of genetic material)

(see item 11)

on Euro  
 summary  
 sheet

same error

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TIME: 11:46:47

Input Set : A:\SEQLIST.TXT

Output Set: N:\CRF4\02022006\J537847.raw

66 <223> OTHER INFORMATION: Artificial peptide sequence

W--> 68 <221> NAME/KEY: PEPTIDE

69 <222> LOCATION: (1)...(9)

70 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding

W--> 72 <400> 3

73 Cys Ser Ser Asn Thr Ala Pro His Cys

74 1 5

77 <210> SEQ ID NO: 4

78 <211> LENGTH: 9

79 <212> TYPE: PRT

80 <213> ORGANISM: Artificial Sequence

82 <220> FEATURE:

83 <223> OTHER INFORMATION: Artificial peptide sequence

W--> 85 <221> NAME/KEY: PEPTIDE

86 <222> LOCATION: (1)...(9)

87 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding

W--> 89 <400> 4

90 Cys His Val Leu Pro Asn Gly Asn Cys

91 1 5

94 <210> SEQ ID NO: 5

95 <211> LENGTH: 10

96 <212> TYPE: PRT

97 <213> ORGANISM: Artificial Sequence

99 <220> FEATURE:

100 <223> OTHER INFORMATION: Artificial peptide sequence

W--> 102 <221> NAME/KEY: PEPTIDE

103 <222> LOCATION: (1)...(9) (10)? since 10 amino acids are in the sequence

104 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding

W--> 106 <400> 5

107 Cys Lys Pro Gly Ile Tyr Pro Ser Leu Cys

108 1 5 10

111 <210> SEQ ID NO: 6

112 <211> LENGTH: 9

113 <212> TYPE: PRT

114 <213> ORGANISM: Artificial Sequence

116 <220> FEATURE:

117 <223> OTHER INFORMATION: Artificial peptide sequence

W--> 119 <221> NAME/KEY: PEPTIDE

120 <222> LOCATION: (1)...(9)

121 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding

W--> 123 <400> 6

124 Cys Gln Thr Ala Arg Thr Pro Ala Cys

125 1 5

128 <210> SEQ ID NO: 7

129 <211> LENGTH: 9

130 <212> TYPE: PRT

131 <213> ORGANISM: Artificial Sequence

133 <220> FEATURE:

134 <223> OTHER INFORMATION: Artificial peptide sequence

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Input Set : A:\SEQLIST.TXT

Output Set: N:\CRF4\02022006\J537847.raw

W--> 136 <221> NAME/KEY: PEPTIDE  
137 <222> LOCATION: (1)...(9)  
138 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding

W--> 140 <400> 7  
141 Cys Asn Gln Ser Gln Pro Lys His Cys  
142 1 5  
145 <210> SEQ ID NO: 8  
146 <211> LENGTH: 9  
147 <212> TYPE: PRT  
148 <213> ORGANISM: Artificial Sequence  
150 <220> FEATURE:  
151 <223> OTHER INFORMATION: Artificial peptide sequence

W--> 153 <221> NAME/KEY: PEPTIDE  
154 <222> LOCATION: (1)...(9)  
155 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding

W--> 157 <400> 8  
158 Cys Thr Pro Ser Lys Ile Ser Val Cys  
159 1 5  
162 <210> SEQ ID NO: 9  
163 <211> LENGTH: 9  
164 <212> TYPE: PRT  
165 <213> ORGANISM: Artificial Sequence  
167 <220> FEATURE:  
168 <223> OTHER INFORMATION: Artificial peptide sequence

W--> 170 <221> NAME/KEY: PEPTIDE  
171 <222> LOCATION: (1)...(9)  
172 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding

W--> 174 <400> 9  
175 Cys Val Ser Pro Gly Pro Arg Leu Cys  
176 1 5  
179 <210> SEQ ID NO: 10  
180 <211> LENGTH: 9  
181 <212> TYPE: PRT  
182 <213> ORGANISM: Artificial Sequence  
184 <220> FEATURE:  
185 <223> OTHER INFORMATION: Artificial peptide sequence

W--> 187 <221> NAME/KEY: PEPTIDE  
188 <222> LOCATION: (1)...(9)  
189 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding

W--> 191 <400> 10  
192 Cys Tyr Ala Leu Ser Gly Val Pro Cys  
193 1 5  
196 <210> SEQ ID NO: 11  
197 <211> LENGTH: 9  
198 <212> TYPE: PRT  
199 <213> ORGANISM: Artificial Sequence  
201 <220> FEATURE:  
202 <223> OTHER INFORMATION: Artificial peptide sequence

W--> 204 <221> NAME/KEY: PEPTIDE

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Input Set : A:\SEQLIST.TXT

Output Set: N:\CRF4\02022006\J537847.raw

205 <222> LOCATION: (1)...(9)  
206 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding  
W--> 208 <400> 11  
209 Cys Lys His Pro Pro Gln Pro Phe Cys  
210 1 5  
213 <210> SEQ ID NO: 12  
214 <211> LENGTH: 9  
215 <212> TYPE: PRT  
216 <213> ORGANISM: Artificial Sequence  
218 <220> FEATURE:  
219 <223> OTHER INFORMATION: Artificial peptide sequence  
W--> 221 <221> NAME/KEY: PEPTIDE  
222 <222> LOCATION: (1)...(9)  
223 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding  
W--> 225 <400> 12  
226 Cys His Gln Ser Lys Pro Leu Leu Cys  
227 1 5  
230 <210> SEQ ID NO: 13  
231 <211> LENGTH: 9  
232 <212> TYPE: PRT  
233 <213> ORGANISM: Artificial Sequence  
235 <220> FEATURE:  
236 <223> OTHER INFORMATION: Artificial peptide sequence  
W--> 238 <221> NAME/KEY: PEPTIDE  
239 <222> LOCATION: (1)...(9)  
240 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding  
W--> 242 <400> 13  
243 Cys Pro Gly Pro Phe Ser Asn Trp Cys  
244 1 5  
247 <210> SEQ ID NO: 14  
248 <211> LENGTH: 9  
249 <212> TYPE: PRT  
250 <213> ORGANISM: Artificial Sequence  
252 <220> FEATURE:  
253 <223> OTHER INFORMATION: Artificial peptide sequence  
W--> 255 <221> NAME/KEY: PEPTIDE  
256 <222> LOCATION: (1)...(9)  
257 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding  
W--> 259 <400> 14  
260 Cys Pro His Lys Thr His Leu Pro Cys  
261 1 5  
264 <210> SEQ ID NO: 15  
265 <211> LENGTH: 9  
266 <212> TYPE: PRT  
267 <213> ORGANISM: Artificial Sequence  
269 <220> FEATURE:  
270 <223> OTHER INFORMATION: Artificial peptide sequence  
W--> 272 <221> NAME/KEY: PEPTIDE  
273 <222> LOCATION: (1)...(9)

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Input Set : A:\SEQLIST.TXT

Output Set: N:\CRF4\02022006\J537847.raw

274 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding  
W--> 276 <400> 15  
277 Cys Val Phe Pro Leu Ser His Tyr Cys  
278 1 5  
281 <210> SEQ ID NO: 16  
282 <211> LENGTH: 9  
283 <212> TYPE: PRT  
284 <213> ORGANISM: Artificial Sequence  
286 <220> FEATURE:  
287 <223> OTHER INFORMATION: Artificial peptide sequence  
W--> 289 <221> NAME/KEY: PEPTIDE  
290 <222> LOCATION: (1)...(9)  
291 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding  
W--> 293 <400> 16  
294 Cys Asn Met Ile Ala Pro Ser Ser Cys  
295 1 5  
298 <210> SEQ ID NO: 17  
299 <211> LENGTH: 9  
300 <212> TYPE: PRT  
301 <213> ORGANISM: Artificial Sequence  
303 <220> FEATURE:  
304 <223> OTHER INFORMATION: Artificial peptide sequence  
W--> 306 <221> NAME/KEY: PEPTIDE  
307 <222> LOCATION: (1)...(9)  
308 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding  
W--> 310 <400> 17  
311 Cys Thr Leu Gly Met Gln Phe Gln Cys  
312 1 5  
315 <210> SEQ ID NO: 18  
316 <211> LENGTH: 9  
317 <212> TYPE: PRT  
318 <213> ORGANISM: Artificial Sequence  
320 <220> FEATURE:  
321 <223> OTHER INFORMATION: Artificial peptide sequence  
W--> 323 <221> NAME/KEY: PEPTIDE  
324 <222> LOCATION: (1)...(9)  
325 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding  
W--> 327 <400> 18  
328 Cys Thr Asn Pro Thr Gly Met Leu Cys  
329 1 5  
332 <210> SEQ ID NO: 19  
333 <211> LENGTH: 9  
334 <212> TYPE: PRT  
335 <213> ORGANISM: Artificial Sequence  
337 <220> FEATURE:  
338 <223> OTHER INFORMATION: Artificial peptide sequence  
W--> 340 <221> NAME/KEY: PEPTIDE  
341 <222> LOCATION: (1)...(9)  
342 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding

Please correct  
this type of error  
in subsequent  
sequences

RAW SEQUENCE LISTING ERROR SUMMARY  
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Input Set : A:\SEQLIST.TXT  
Output Set: N:\CRF4\02022006\J537847.raw

*FYI*  
Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:43; Xaa Pos. 2,3,8.



## VERIFICATION SUMMARY

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Input Set : A:\SEQLIST.TXT

Output Set: N:\CRF4\02022006\J537847.raw

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:34 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:38 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1  
L:51 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:55 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:68 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:72 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3  
L:85 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:89 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4  
L:102 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:106 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5  
L:119 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:123 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6  
L:136 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:140 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7  
L:153 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:157 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8  
L:170 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:174 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9  
L:187 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:191 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10  
L:204 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:208 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11  
L:221 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:225 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12  
L:238 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:242 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:13  
L:255 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:259 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14  
L:272 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:276 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:15  
L:289 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:293 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16  
L:306 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:310 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:17  
L:323 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:327 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18  
L:340 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:344 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19  
L:357 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:361 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:20  
L:374 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:378 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:21  
L:391 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:395 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22  
L:408 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:412 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23  
L:425 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!

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Input Set : A:\SEQLIST.TXT

Output Set: N:\CRF4\02022006\J537847.raw

L:429 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24  
L:442 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:446 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25  
L:459 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:463 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26  
L:476 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:480 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:27  
L:493 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:497 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:28  
L:510 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:514 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:29  
L:527 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:531 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:30  
L:544 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:548 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:31  
L:561 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:565 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:32  
L:578 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:582 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:33  
L:595 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:599 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:34  
L:612 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:616 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:35  
L:629 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:633 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36  
L:646 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:650 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:37  
L:663 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:668 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:38  
L:679 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:684 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:39  
L:695 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:701 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:40  
L:712 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:718 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:41  
L:729 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:733 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:42  
L:746 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:750 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:43  
L:751 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:0